

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/806,062

Source: _____

Date Processed by STIC: _____

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RAW SEQUENCE LISTING

DATE: 06/06/2005

PATENT APPLICATION: US/10/806,062

TIME: 07:58:32

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Output Set: N:\CRF4\06062005\J806062.raw

1 <110> APPLICANT: EDWARDS, ALED
2 DHARAMSI, AKIL
3 VEDADI, MASOUD
4 ALAM, MUHAMMAD ZAHOR
5 DOMAGALA, MEGAN
6 HOUSTON, SIMON
7 LAM, ROBERT
8 LI, QIN
9 NETHERY, KATHLEEN
10 NG, IVY
11 PINDER, BENJAMIN
12 VIOLA, CRISTINA
13 WREZEL, OLGA
14 KANAGARAJAH, DHUSHY
15 MANSOURY, KAMRAN
16 NECAKOV, ALEKSANDAR SASHA
17 VALLEE, FRANCOIS
18 MCDONALD, MERRY-LYNN
19 <120> TITLE OF INVENTION: NOVEL PURIFIED POLYPEPTIDES FROM BACTERIA
20 <130> FILE REFERENCE: IPT-101.01
21 <140> CURRENT APPLICATION NUMBER: US/10/806,062
22 <141> CURRENT FILING DATE: 2004-03-22
23 <150> PRIOR APPLICATION NUMBER: PCT/CA02/01428
24 <151> PRIOR FILING DATE: 2002-09-20
25 <150> PRIOR APPLICATION NUMBER: 60/324,152
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38 <151> PRIOR FILING DATE: 2001-11-27
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40 <151> PRIOR FILING DATE: 2001-12-18
41 <150> PRIOR APPLICATION NUMBER: 60/341,776
42 <151> PRIOR FILING DATE: 2001-12-18
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Input Set : N:\CrF4\Refhold\10_folder\J806062MOD.raw

Output Set: N:\CRF4\06062005\J806062.raw

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162     gaagtaaagc ctacagatta tttagttaca gagttgtcgt cattccagtt acttggaatc 540
163     gaaaagtata aaccacacat tgctataatt actaacattt attcggcgca tctagattac 600
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188         35             40             45
189     Ser Gln Asp Ala His Ala Lys Asp Leu Glu Ser Met Gly Ile Ser Val
190         50             55             60
191     Val Ser Gly Ser His Pro Leu Thr Leu Leu Asp Asn Asn Pro Ile Ile
192         65             70             75             80
193     Val Lys Asn Pro Gly Ile Pro Tyr Thr Val Ser Ile Ile Asp Glu Ala
194         85             90             95
195     Val Lys Arg Gly Leu Lys Ile Leu Thr Glu Val Glu Leu Ser Tyr Leu

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198          115          120          125
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200          130          135          140
201 Gly Arg Leu Ser Gly Asn Ile Gly Tyr Val Ala Ser Lys Val Ala Gln
202          145          150          155          160
203 Glu Val Lys Pro Thr Asp Tyr Leu Val Thr Glu Leu Ser Ser Phe Gln
204          165          170          175
205 Leu Leu Gly Ile Glu Lys Tyr Lys Pro His Ile Ala Ile Ile Thr Asn
206          180          185          190
207 Ile Tyr Ser Ala His Leu Asp Tyr His Glu Asn Leu Glu Asn Tyr Gln
208          195          200          205
209 Asn Ala Lys Lys Gln Ile Tyr Lys Asn Gln Thr Glu Glu Asp Tyr Leu
210          210          215          220
211 Ile Cys Asn Tyr His Gln Arg Gln Val Ile Glu Ser Glu Glu Leu Lys
212          225          230          235          240
213 Ala Lys Thr Leu Tyr Phe Ser Thr Gln Gln Glu Val Asp Gly Ile Tyr
214          245          250          255
215 Ile Lys Asp Gly Phe Ile Val Tyr Lys Gly Val Arg Ile Ile Asn Thr
216          260          265          270
217 Glu Asp Leu Val Leu Pro Gly Glu His Asn Leu Glu Asn Ile Leu Ala
218          275          280          285
219 Ala Val Leu Ala Cys Ile Leu Ala Gly Val Pro Ile Lys Ala Ile Ile
220          290          295          300
221 Asp Ser Leu Thr Thr Phe Ser Gly Ile Glu His Arg Leu Gln Tyr Val
222          305          310          315          320
223 Gly Thr Asn Arg Thr Asn Lys Tyr Tyr Asn Asp Ser Lys Ala Thr Asn
224          325          330          335
225 Thr Leu Ala Thr Gln Phe Ala Leu Asn Ser Phe Asn Gln Pro Ile Ile
226          340          345          350
227 Trp Leu Cys Gly Gly Leu Asp Arg Gly Asn Glu Phe Asp Glu Leu Ile
228          355          360          365
229 Pro Tyr Met Glu Asn Val Arg Ala Met Val Val Phe Gly Gln Thr Lys
230          370          375          380
231 Ala Lys Phe Ala Lys Leu Gly Asn Ser Gln Gly Lys Ser Val Ile Glu
232          385          390          395          400
233 Ala Asn Asn Val Glu Asp Ala Val Asp Lys Val Gln Asp Ile Ile Glu
234          405          410          415
235 Pro Asn Asp Val Val Leu Leu Ser Pro Ala Cys Ala Ser Trp Asp Gln
236          420          425          430
237 Tyr Ser Thr Phe Glu Glu Arg Gly Glu Lys Phe Ile Glu Arg Phe Arg
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239 Ala His Leu Pro Ser Tyr
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243 <211> LENGTH: 1365
244 <212> TYPE: DNA
245 <213> ORGANISM: Staphylococcus aureus

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:323; Xaa Pos. 3
Seq#:324; Xaa Pos. 2,5

VERIFICATION SUMMARY

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Input Set : N:\Crf4\Refhold\10_folder\J806062MOD.raw

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L:6371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:323 after pos.:0

L:6390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:324 after pos.:0